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
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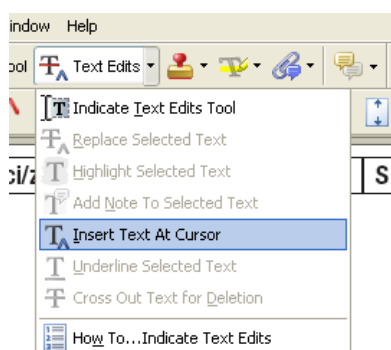
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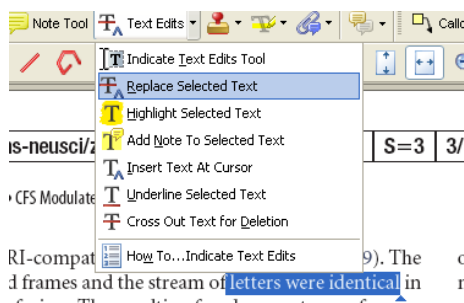
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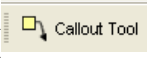
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Subject	Target contrast (%)
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S3	15
S4	20
Mean $\pm$ SEM	14.75 $\pm$ 1.89

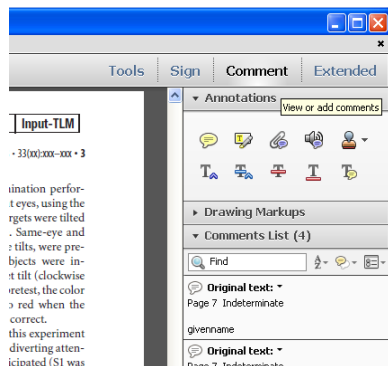
Each row corresponds to a different subject. Bottom row, mean and SEM across performance for target and mask presented to different eyes; well above chance level.

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# Amyloid $\beta$ Oligomers Disrupt Blood–CSF Barrier Integrity by Activating Matrix Metalloproteinases

AQ: au  
AQ: or

Marjana Brkic,<sup>1,2,5\*</sup>  Sriram Balusu,<sup>1,2\*</sup> Elien Van Wonterghem,<sup>1,2</sup> Nina Gorlé,<sup>1,2</sup> Iryna Benilova,<sup>3,4</sup> Anna Kremer,<sup>1,2</sup> Inge Van Hove,<sup>6</sup> Lieve Moons,<sup>6</sup> Bart De Strooper,<sup>3,4</sup> Selma Kanazir,<sup>5</sup> Claude Libert,<sup>1,2†</sup> and  Roosmarijn E. Vandenbroucke<sup>1,2†</sup>

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AQ: A

The blood–CSF barrier (BCSFB) consists of a monolayer of choroid plexus epithelial (CPE) cells that maintain CNS homeostasis by producing CSF and restricting the passage of undesirable molecules and pathogens into the brain. Alzheimer’s disease is the most common progressive neurodegenerative disorder and is characterized by the presence of amyloid  $\beta$  (A $\beta$ ) plaques and neurofibrillary tangles in the brain. Recent research shows that Alzheimer’s disease is associated with morphological changes in CPE cells and compromised production of CSF. Here, we studied the direct effects of A $\beta$  on the functionality of the BCSFB. Intracerebroventricular injection of A $\beta$ 1–42 oligomers into the cerebral ventricles of mice, a validated Alzheimer’s disease model, caused induction of a cascade of detrimental events, including increased inflammatory gene expression in CPE cells and increased levels of proinflammatory cytokines and chemokines in the CSF. It also rapidly affected CPE cell morphology and tight junction protein levels. These changes were associated with loss of BCSFB integrity, as shown by an increase in BCSFB leakage. A $\beta$ 1–42 oligomers also increased matrix metalloproteinase (MMP) gene expression in the CPE and its activity in CSF. Interestingly, BCSFB disruption induced by A $\beta$ 1–42 oligomers did not occur in the presence of a broad-spectrum MMP inhibitor or in MMP3-deficient mice. These data provide evidence that MMPs are essential for the BCSFB leakage induced by A $\beta$ 1–42 oligomers. Our results reveal that Alzheimer’s disease-associated soluble A $\beta$ 1–42 oligomers induce BCSFB dysfunction and suggest MMPs as a possible therapeutic target.

AQ: B

**Key words:** alzheimer’s disease; amyloid  $\beta$  toxicity; blood–CSF barrier; choroid plexus; matrix metalloproteinases

## Significance Statement

No treatments are yet available to cure Alzheimer’s disease; however, soluble A $\beta$  oligomers are believed to play a crucial role in the neuroinflammation that is observed in this disease. Here, we studied the effect of A $\beta$  oligomers on the often neglected barrier between blood and brain, called the blood–CSF barrier (BCSFB). This BCSFB is formed by the choroid plexus epithelial cells and is important in maintaining brain homeostasis. We observed A $\beta$  oligomer-induced changes in morphology and loss of BCSFB integrity that might play a role in Alzheimer’s disease progression. Strikingly, both inhibition of matrix metalloproteinase (MMP) activity and MMP3 deficiency could protect against the detrimental effects of A $\beta$  oligomer. Clearly, our results suggest that MMP inhibition might have therapeutic potential.

## Introduction

Fn1

Alzheimer’s disease is the most common progressive form of dementia, characterized by synaptic loss, neurodegeneration, and

impairment of cognitive function. Overproduction and decreased clearance of amyloid  $\beta$  (A $\beta$ ) peptide, followed by formation of amyloid plaques, is believed to be the central player of Alzheimer’s disease (Deane et al., 2009; Alvira-Botero and Carro, 2010). More than a decade ago, it was observed that progression

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The authors declare no competing financial interests.

\*M.B. and S.B. contributed equally to this work.



of the disease is more strongly correlated with the presence of soluble A $\beta$  than with the number of plaques (McLean et al., 1999). This A $\beta$  molecular form, which is an intermediate between monomeric A $\beta$  and insoluble amyloid plaques, is now recognized as important mediator in the pathology of Alzheimer's disease. Recently, it was shown that mice injected with oligomerized A $\beta$ 1–42 display neuronal cell loss, tau hyperphosphorylation, and impairment of hippocampus-dependent memory (Brouillette et al., 2012; Ledo et al., 2013). Similarly, injection of A $\beta$ 1–42 in the brain ventricles of zebrafish embryos leads to cognitive deficits and tau hyperphosphorylation (Nery et al., 2014).

The microenvironment in the brain is strictly maintained by two main barriers: the blood–brain barrier (BBB) in the brain parenchyma and the blood–CSF barrier (BCSFB) in the choroid plexus (CP). The BBB separates brain interstitial fluid from blood at the level of brain capillaries, which are formed by endothelial cells connected through tight junctions, restricting paracellular transport. In addition, pericytes and astrocytic end feet contribute to strict insulation of the BBB (Zlokovic, 2008). In contrast, capillaries in the CP are highly fenestrated due to the absence of tight junctions. Therefore, the BCSFB is formed by CP epithelial (CPE) cells that are firmly interconnected by tight junctions separating blood from CSF. CPE cells also contain many specific transport systems and receptors that provide them with an active role in the regulation of transport from blood to CSF and vice versa. In addition, the BCSFB acts as a relay station that senses inflammation signals from both the CNS (Batra et al., 2010; Sharma et al., 2010; Simard et al., 2011) and the periphery (Marques et al., 2007; Marques et al., 2009a; Marques et al., 2009b; Vandenbroucke et al., 2012). CPE cells respond to inflammatory stimuli by producing proinflammatory molecules, which is often associated with disturbance of barrier integrity and results in transmission of the inflammatory signals to the rest of the brain (Mitchell et al., 2009; Coisne and Engelhardt, 2011) and increased white blood cell influx into the brain (Demeestere et al., 2015).

Alzheimer's disease is associated with numerous changes in CP morphology and function, of which the most prominent are decreased CSF production, changes in metabolic activity, and reduced clearance of toxins, including A $\beta$  (Serot et al., 2000; Serot et al., 2003; Emerich et al., 2005; Krzyzanowska and Carro, 2012; Serot et al., 2012; Marques et al., 2013; Spector and Johanson, 2013). In addition, the vascular system of the brain is compromised (Zlokovic, 2011). Recent studies suggest that one of the possible causes of BBB disruption in Alzheimer's disease is degeneration and loss of pericytes (Sagare et al., 2013; Winkler et al., 2014). Breakdown of the endothelial and epithelial barriers has been linked to increased activity of matrix metalloproteinases (MMPs), endopeptidases that are implicated in several inflammatory processes (Vandenbroucke and Libert, 2014) and are known to affect tight junction functionality and extracellular matrix composition, eventually aggravating brain inflammation (Zeni et al., 2007; Batra et al., 2010; Vandenbroucke et al., 2012). Several studies have suggested that MMPs play a role in Alzheimer's disease pathogenesis (Wang et al., 2014). For example, CSF and brain MMP levels are higher in Alzheimer's disease patients

than in age-matched controls and MMPs have been implicated in degradation of A $\beta$  (Guo et al., 2006; Yan et al., 2006; Yin et al., 2006; Hanzel et al., 2014; Kauwe et al., 2014; Mroczko et al., 2014). Here, we studied the direct toxic effects of A $\beta$ 1–42 oligomers on the functionality of the CPE.

## Materials and Methods

**Mice.** Female C57BL/6 mice (8–10 weeks old) were purchased from Janvier and housed in our specific pathogen-free (SPF) animal facility. MMP3-deficient mice (Mudgett et al., 1998) in the C57BL/6 background were bred in our SPF facility (MMP3<sup>−/−</sup>). Mice were housed in groups of 4–6/cage with *ad libitum* access to food and water and a 14 h light/10 h dark cycle. All experiments were approved by the ethics committee of the Faculty of Sciences of Ghent University.

**Preparation of A $\beta$ 1–42 oligomers.** Oligomerized A $\beta$ 1–42 was prepared as described previously (Kuperstein et al., 2010; Brouillette et al., 2012). Briefly, A $\beta$ 1–42 (rPeptide; A-1163-1) or scrambled A $\beta$ 1–42 (rPeptide; A-1004-1) was dissolved at 1 mg/ml in hexafluoroisopropanol (HFIP; Sigma-Aldrich; catalog #105228), followed by HFIP removal in a Speed-Vac vacuum concentrator. The resulting peptide film was resolved at 1 mg/ml in DMSO (Sigma-Aldrich; catalog #D4540). Next, the peptide was purified from DMSO on a 5 ml HiTrap desalting column (GE Healthcare; catalog #17-408-01) and eluted with Tris-EDTA buffer (50 mM Tris and 1 mM EDTA, pH 7.5). The resulting peptide concentration was determined using the Thermo Scientific–Pierce Micro BCA Protein Assay (catalog #23225) according to the manufacturer's instructions. Finally, the eluted peptide was allowed to aggregate for 2 h at room temperature and then diluted to 1  $\mu$ g/ml in Tris-EDTA buffer. To confirm the oligomer formation, we characterized the freshly prepared A $\beta$ 1–42 oligomers by analyzing Thioflavin T incorporation into aggregates, performing dot blot analysis, atomic force microscopy analysis, Coomassie-stained SDS-PAGE gel analysis, and cell toxicity assays as described previously (data not shown; Brouillette et al., 2012).

**Injection of A $\beta$ 1–42 oligomers and scrambled peptide in the cerebral ventricles.** Animals were anesthetized with isoflurane and placed in a stereotactic frame. Body temperature was maintained at 37°C using a heating pad. Injection coordinates were measured from the bregma (anterior–posterior −0.07, mediolateral 0.1, dorsoventral −0.3) and were determined using the Franklin and Paxinos mouse brain atlas. A volume of 5  $\mu$ l (1  $\mu$ g/ml peptide) was injected in the right lateral cerebral ventricle using a Hamilton needle. To assess the role of MMPs, we used three treatments: A $\beta$ 1–42 oligomers (in Tris-EDTA buffer) combined with 1  $\mu$ g of broad spectrum MMP inhibitor (GM6001; Merck; CC1100) dissolved in DMSO, A $\beta$ 1–42 oligomers (in Tris-EDTA buffer) combined with DMSO and vehicle for the control group.

**BCSFB and BBB permeability.** BCSFB and BBB permeability were determined as described previously (Vandenbroucke et al., 2012). In brief, 4 kDa FITC-dextran (Sigma-Aldrich; catalog #46944) was injected intravenously 1 h before CSF collection. Two and 6 h after A $\beta$ 1–42 oligomer injection, mice were sedated with ketamine/xylazine and CSF was obtained from the fourth ventricle using the cisterna magna puncture method (Liu and Duff, 2008). Next, mice were transcardially perfused with D-PBS/heparin (0.2% heparin) and brain tissue was isolated. CSF samples were diluted 100-fold in sterile D-PBS and BCSFB leakage was determined by measurement of fluorescence at  $\lambda_{ex}/\lambda_{em}$  = 488/520 nm. Brain samples were cut into small pieces, incubated overnight at 37°C in formamide while shaking, and supernatant was collected after centrifugation for 15 min at maximal speed. Brain fluid was diluted twofold in sterile D-PBS and BBB leakage was determined by measurement of fluorescence at  $\lambda_{ex}/\lambda_{em}$  = 488/520 nm.

**Tissue isolation.** For RNA and protein analysis, mice were transcardially perfused with D-PBS/heparin (0.2% heparin) supplemented with 0.5% bromophenol blue. Brain tissue was dissected out, CP was obtained from all four ventricles, and hippocampus was isolated. Isolated hippocampus was kept in RNAlater (Ambion; catalog #AM7020) and CP was snap frozen in liquid nitrogen. For immunohistochemical analysis, brain was immediately frozen in cryoprotectant (Thermo Scientific; catalog #4583) and stored at −80°C for cryosectioning or fixed in 4% para-

f.L. and R.E.V. contributed equally to this work.

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AQ:C-D

formaldehyde (PFA) followed by paraffin embedding for paraffin sections.

**Immunohistochemistry.** For Occludin and Claudin-5 immunostaining of cryosections, 30  $\mu\text{m}$  sagittal sections were cut using a cryostat (Micron; catalog #HM500) and mounted on slides. After air drying for 2 h, sections were fixed with 1% PFA for 10 min, washed for 5 min in PBS, and permeabilized for 10 min with 0.1% NP-40. After 2 washes with PBS, samples were blocked for 1 h at room temperature with 5% BSA (Sigma-Aldrich; catalog #A2153) and then incubated with primary antibody (Life Technologies; Occl, 1:100, 33–1500; Cldn5, 1:50, 35–2500). After incubation overnight at 4°C, they were washed and incubated with secondary antibody (Thermo Scientific; goat anti-mouse-DyLight633 and goat anti-rabbit-DyLight633, 1:400) diluted in 5% BSA for 90 min at room temperature (RT). For E-cadherin and Iba1 staining, paraffin-embedded brains were sectioned at 4  $\mu\text{m}$  in sagittal orientation, dewaxed, and rinsed in water and PBS before staining. Antigen retrieval was done using citrate buffer (Dako; catalog #S2031), followed by washing in PBS. For E-cadherin staining, slides were incubated overnight at 4°C with 0.01 M  $\text{NaBH}_4$  (Sigma-Aldrich; catalog #452882) to further reduce autofluorescence. After rinsing in PBS, permeabilization was done with 0.05% Tween-20 (Sigma-Aldrich; catalog #P1379) for 30 min at RT. Samples were blocked with 2% BSA for 30 min at RT, followed by incubation for 90 min at RT with primary (E-Cadherin, 1:500; BD Transduction Laboratories; catalog #610181) and secondary antibodies (; Alexa Fluor 568 goat anti-mouse, 1:500; Life Technologies) diluted in 5% BSA. For Iba-1 staining, antigen retrieval was followed by overnight incubation at 4°C with primary antibody (; Iba1, 1:1000; Wako; catalog #019 19741). The next day, slides were washed and incubated with LSAB2 System HRP (DAKO; catalog #K0672) and visualization was done using DAB chromogen. Finally, slides were dehydrated and xylene-based mounting medium was applied. Cells were visualized using Olympus BX51 microscope. Iba1-positive cells were counted in predefined area of the brain, including both cortex and hippocampus, using Fiji (<http://fiji.sc/Fiji>). Microglia were classified into resting and activated according to adopted criteria (Hains and Waxman, 2006).

**Cytokine/chemokine measurements.** Cytokines and chemokines in CSF were measured using the Bio-Plex cytokine assays (Bio-Rad; catalog #M60009RDPD) according to the manufacturer's instructions.

**RNA isolation.** Total RNA was isolated with the (mi)RNeasy kit (Qiagen; catalog #74106). RNA concentration and purity were determined spectrophotometrically using the Nanodrop Technologies ND-1000.

**Real-time qPCR.** After RNA isolation, cDNA was synthesized by using a cDNA Synthesis Kit (Bio-Rad; catalog #172-5038). Real-time qPCR was performed on the Light Cycler 480 system (Roche) using the Light-Cycler 480 SYBR Green I Master mix (Roche; catalog #04887352001) or the SensiFAST SYBR No-ROX Kit (Bioline; catalog #BIO-98002). Expression levels were normalized to the expression of the two or three most stable reference genes, as determined by the geNorm Housekeeping Gene Selection Software (Vandesompele et al., 2002): *Hprt*, *Rpl*, and *Ubc* for CP and *Gapdh*, *Rpl*, and *Ubc* for hippocampus. The sequences of the forward and reverse primers for the different genes are provided in Table 1.

**MMP activity.** MMP activity in the CSF was analyzed using the Omni-MMP fluorogenic substrate kit according to manufacturer's guidelines (Enzo Life Sciences; catalog #BML-AK016). In brief, CSF samples were diluted 50-fold in assay buffer and MMP activity was determined by measuring the increase in fluorescence at  $\lambda_{\text{ex}}/\lambda_{\text{em}} = 320/460 \text{ nm}$ .

**Western blot.** For Occludin Western blot analysis, CP samples from two mice were pooled and homogenized in 50  $\mu\text{l}$  of lysis buffer containing 0.5% CHAPS (Sigma-Aldrich; catalog #C9426) and protease inhibitor complete tablet (Roche Applied Science; catalog #11 873 580 001). Protein concentration was determined using the Thermo Scientific–Pierce Micro BCA Protein Assay and 60  $\mu\text{g}$  of proteins were loaded on a 12.5% SDS-PAGE gel. For MMP3 Western blot analysis, 2  $\mu\text{l}$  of each sample, derived from scrambled and  $\text{A}\beta$ 1–42 oligomer injected mice, was loaded on a 12.5% SDS-PAGE gel. Further protocol remained the same for both Occludin and MMP3 detection. Semidry transfer was performed for 1 h at constant current (45 mA) and membranes were

**Table 1. Overview of the sequences of the forward and reverse primers used for qPCR analysis**

Gene	Forward	Reverse
<i>Il1<math>\beta</math></i>	CACCTCACAAGCAGACACAAG	GCATTAGAAACAGTCAGCCCATAC
<i>Il6</i>	TAGTCCTTCTACCCCAATTTCC	TTGGTCCTTAGCAGCTCCTTC
<i>Tnf</i>	ACCCTGGTATGAGCCATATAC	ACACCCATTCCTTCACAGAG
<i>Ocln</i>	CCAGGCAGCGTGTCTCT	TTCTAATAACAGTCACCTGAGGCG
<i>Cldn5</i>	GCAAGGTGTATGAATCTGTGCT	GTCAGGTAACAAGAGTGCCCA
<i>Zo1</i>	AGGACCAAAAGCATGTGAG	GGCATTCTGCTGTTTACA
<i>Zo3</i>	ACCCTATGGCTGGGCTTC	CCCGGGTACAACGTGTCC
<i>Cldn1</i>	TCTACGAGGAGCTGTGGATG	TCAGATTACAGCAAGGAGTCG
<i>Ecdh</i>	TCGGAAGACTCCCGATTCAA	CGGACGAGGAACCTGGTCTC
<i>Mmp1a</i>	CCTGTATGAGACGTGGACCA	ATGTGGTGTGTTCACCTGT
<i>Mmp2</i>	AGATCTTCTTCAAGGACCGGT	GGCTGTGTCAGTGGCTGGGTA
<i>Mmp3</i>	AGTCTACAAGCTCCACAG	TTGGTATGTCTCAGGTTC
<i>Mmp8</i>	ATTCACAGGAGTGCCAAGC	TGATTGTCATATCCAGCACTGG
<i>Mmp9</i>	CTGGACAGCCAGACACTAAG	CTCGCGGAAGTCTTCAGAG
<i>Mmp13</i>	TTTATTGTTGCTGCCATGA	GGTCTTGGAGTATCCAGA
<i>Mmp14</i>	CAGTATGGCTACCTACCTCCAG	GCCTTGGCTGCTACTGTAAA
<i>Rpl</i>	CCTGCTGCTCAAGGTT	TGGTGTGCTGCTGCTACTT
<i>Ubc</i>	AGGTCAAACAGGAAGACAGCGTA	TCACACCAAGAACAAGCACA
<i>Gapdh</i>	TGAAGCAGGCATCTGAGGG	CGAAGGTGGAAGAGTGGGAG
<i>Hprt</i>	AGTGTGGATACAGGCCAGAC	CGTGATTCAATCCCTGAAGT
$\beta$ 2M	ATGCACGAGAAAGAAATAGCAA	AGCTATCTAGGATATTCCAATTTTGA
$\beta$ act	GCTCTAGGCGGACTGTACTGA	GCCATGCCAATGTTGCTCTTAT

blocked for 1 h using Odyssey blocking buffer (Li-Cor; catalog #927-40000), followed by overnight incubation with primary antibody at 4°C (anti-Occludin, 1:250, Life Technologies, catalog #33-1500; anti-MMP3, 1:100, Abcam, catalog #ac52915). Next, membranes were incubated with the secondary antibody (Thermo Scientific anti-mouse DyLight 680, 1:1000) for 2 h at RT and bands were visualized using Odyssey software.

**Serial block-face scanning electron microscopy.** For serial block-face scanning electron microscopy (SBF-SEM), CP tissue was dissected and immediately transferred into fixation buffer (2% paraformaldehyde, Sigma-Aldrich; 2.5% glutaraldehyde, Electron Microscopy Sciences in 0.15 M cacodylate buffer, pH 7.4). After overnight fixation at 4°C, samples were washed 3  $\times$  5 min in cacodylate buffer and subsequently osmicated in 2% osmium (EMS), 1.5% ferrocyanide, and 2 mM  $\text{CaCl}_2$  in cacodylate buffer for 1 h on ice, and then washed extensively in ultrapure water (UPW). This was followed by incubation in 1% thiocarbonyhydrazide (20 min), washes in UPW, and a second osmication in 2% osmium in UPW (30 min). The samples were washed 5  $\times$  3 min in UPW and placed in 2% uranyl acetate at 4°C overnight. The following day, they were stained with Walton's lead aspartate stain for 30 min at 60°C. For this, a 30 mM L-aspartic acid solution was used to freshly dissolve lead nitrate (final concentration 20 mM, pH 5.5). The solution was filtered after incubation for 30 min at 60°C.

After the final washes, the samples were dehydrated using a series of ice-cold solutions of increasing ethanol concentration (30%, 50%, 70%, 90%, and twice 100%), followed by two dehydrations of 30 min in 100% acetone. Subsequent infiltration with resin (Durcupan; EMS) was done by first incubating the samples in 50% resin in acetone for 4 h, followed by at least 5 changes of fresh 100% resin (including 2 overnight incubations). Next, samples were embedded in fresh resin and cured in the oven at 65°C for 72 h.

For SBF-SEM, the resin-embedded samples were mounted on an aluminum specimen pin (Gatan) using conductive epoxy (Circuit Works). The specimens were trimmed in a pyramid shape using an ultramicrotome (Ultracut; Leica) and the block surface was trimmed until smooth and at least a small part of tissue was present at the block face. Next, samples were coated with 5 nm Pt in a Quorum Q 150T ES sputter coater (Quorum Technologies). The aluminum pins were placed in the Gatan 3View2XP in a Zeiss Merlin SEM for imaging at 1.6 kV with a Gatan Digiscan II ESB detector. The Gatan 3view2XP was set to section 300 sections of 70 nm. IMOD (<http://bio3d.colorado.edu/imod/>) and Fiji (Schindelin et al., 2012) were used for registration of the 3D image



stack and conversion to TIFF file format. Representation of the cell in 3D movies and snapshots was done in Imaris (BitPlane). 3D modeling of CPE cells was performed using IMOD software.

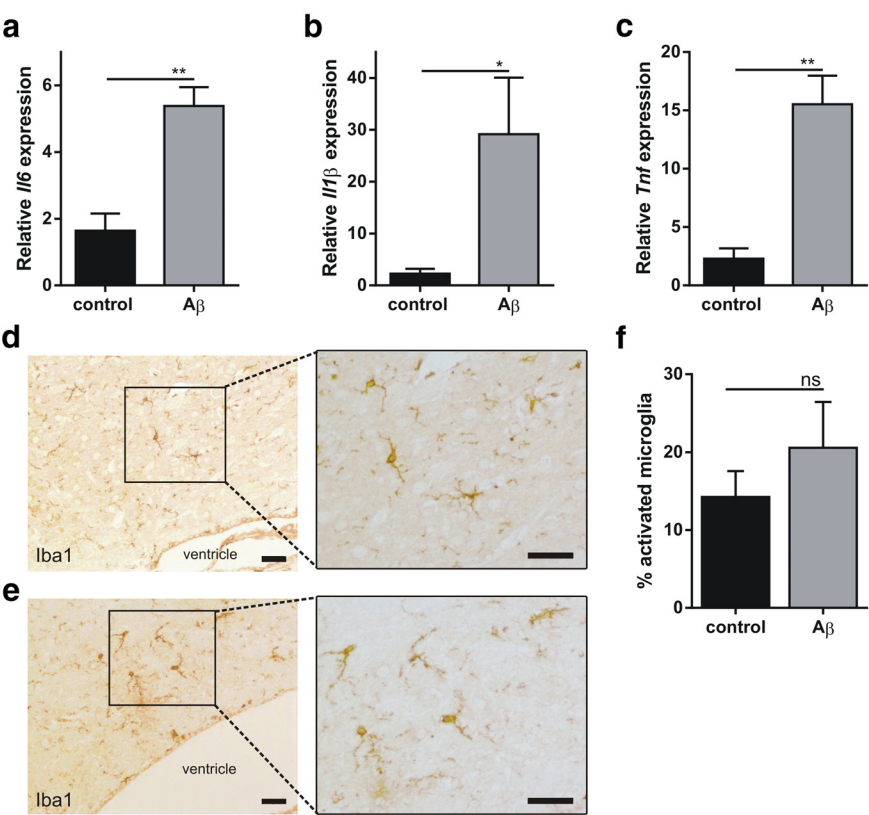
**Statistics.** Data were analyzed using Graph-Pad Prism and Student's *t* test and are presented as means  $\pm$  SEM. Significance levels are indicated on the graphs as follows: \*0.01  $\leq p < 0.05$ ; \*\*0.001  $\leq p < 0.01$ ; \*\*\*0.0001  $\leq p < 0.001$ ; and \*\*\*\**p* < 0.0001.

Results

Intracerebroventricular administration of A $\beta$ 1–42 oligomers induces inflammation in brain, CP, and CSF

In Alzheimer's disease, A $\beta$  and its deposits, along with neurofibrillary tangles, induce a chronic local upregulation of inflammatory mediators (Akiyama et al., 2000). We prepared A $\beta$ 1–42 oligomers by resuspending A $\beta$ 1–42 peptide film in Tris-EDTA buffer as described previously (Kuperstein et al., 2010), followed by incubation for 2 h to allow oligomerization. Next, C57BL/6 mice were injected intracerebroventricularly with A $\beta$ 1–42 oligomers (Brouillette et al., 2012) and the effect on brain inflammation was studied. Gene expression analysis of the hippocampus revealed that several inflammatory genes were significantly upregulated 6 h after A $\beta$ 1–42 oligomer injection: *Il6*, *Il1 $\beta$* , and *Tnf* levels are shown in Figure 1*a–c*, respectively. In contrast, Iba1 immunostaining did not reveal a significant increase in microglia activation in mice 6 h after intracerebroventricular injection of A $\beta$ 1–42 (Fig. 1*e,f*) compared with injection of scrambled peptide (Fig. 1*d,f*). The CP is one of the sites involved in the production, deposition, and clearance of A $\beta$  in the brain (Crossgrove et al., 2005; Crossgrove et al., 2007; Krzyzanowska and Carro, 2012). To study the direct effect of A $\beta$ 1–42 oligomers on inflammation in the CP and hippocampus, CSF, CP, and hippocampus were isolated 2 and 6 h after intracerebroventricular injection of scrambled peptide or A $\beta$ 1–42 oligomers and total RNA was extracted from CP and hippocampus. Next, we determined mRNA levels of several proinflammatory cytokines (Fig. 2*a–h*). *Il1 $\beta$* , *Il6*, *Tnf*, and *Inos* were significantly upregulated 2 h after intracerebroventricular injection of A $\beta$ 1–42 compared with scrambled peptide. The upregulation was even more pronounced 6 h after intracerebroventricular injection in CP (Fig. 2*a–d*). Similarly, we determined mRNA levels of the proinflammatory cytokines in the hippocampus (Fig. 2*e–h*). As observed in the CP, there was a significant increase in the *Il1 $\beta$* , *Il6*, *Tnf*, and *Inos* mRNA expression at 2 h. With the exception of *iNos*, these effects were more pronounced at 6 h.

Next, to determine whether this upregulation in cytokine expression was accompanied by inflammatory protein expression, we evaluated cytokine levels in CSF using the BioPlex assay (Fig. 2*i–l*). The significantly increased presence of the proinflammatory cytokines IL-1 $\beta$ , IL-6, TNF $\alpha$ , and IFN $\gamma$  in CSF at both time points correlated with the elevated mRNA levels observed in the CP: they were significantly increased 2 and 6 h after intracerebroventricular injection of A $\beta$ 1–42. In parallel, we measured several chemokines in the CSF: MIP-1 $\alpha$ , MIP-1 $\beta$ , MCP-1, and GM-CSF



**Figure 1.** Analysis of inflammation in the hippocampus after intracerebroventricular injection of oligomerized A $\beta$ 1–42 in the cerebral ventricles. *a–c*, mRNA expression analysis of *Il1 $\beta$*  (*a*), *Il6* (*b*), and *Tnf* (*c*) in the hippocampus 6 h after intracerebroventricular injection of A $\beta$ 1–42 in C57BL/6 mice compared with control hippocampus samples (*n* = 4). *d, e*, Representative images of Iba1 staining of brain sections from C57BL/6 mice 6 h after intracerebroventricular injection with scrambled peptide (*d*) or A $\beta$ 1–42 oligomers (*e*). As indicated on the images, the region surrounding the lateral ventricle is represented. *f*, Quantification of the percentage of activated microglia in cortex and hippocampus of scrambled and A $\beta$ 1–42 oligomer injected mice (*n* = 5). Scale bar, 25  $\mu$ m.

were all upregulated at the tested time points in the presence of A $\beta$ 1–42 (Fig. 2*m–p*).

Our data show that A $\beta$ 1–42 oligomers induce a strong inflammatory response in the brain that is more pronounced in the CP than in the rest of the brain. This is reflected by increased cytokine and chemokine levels in the CSF.

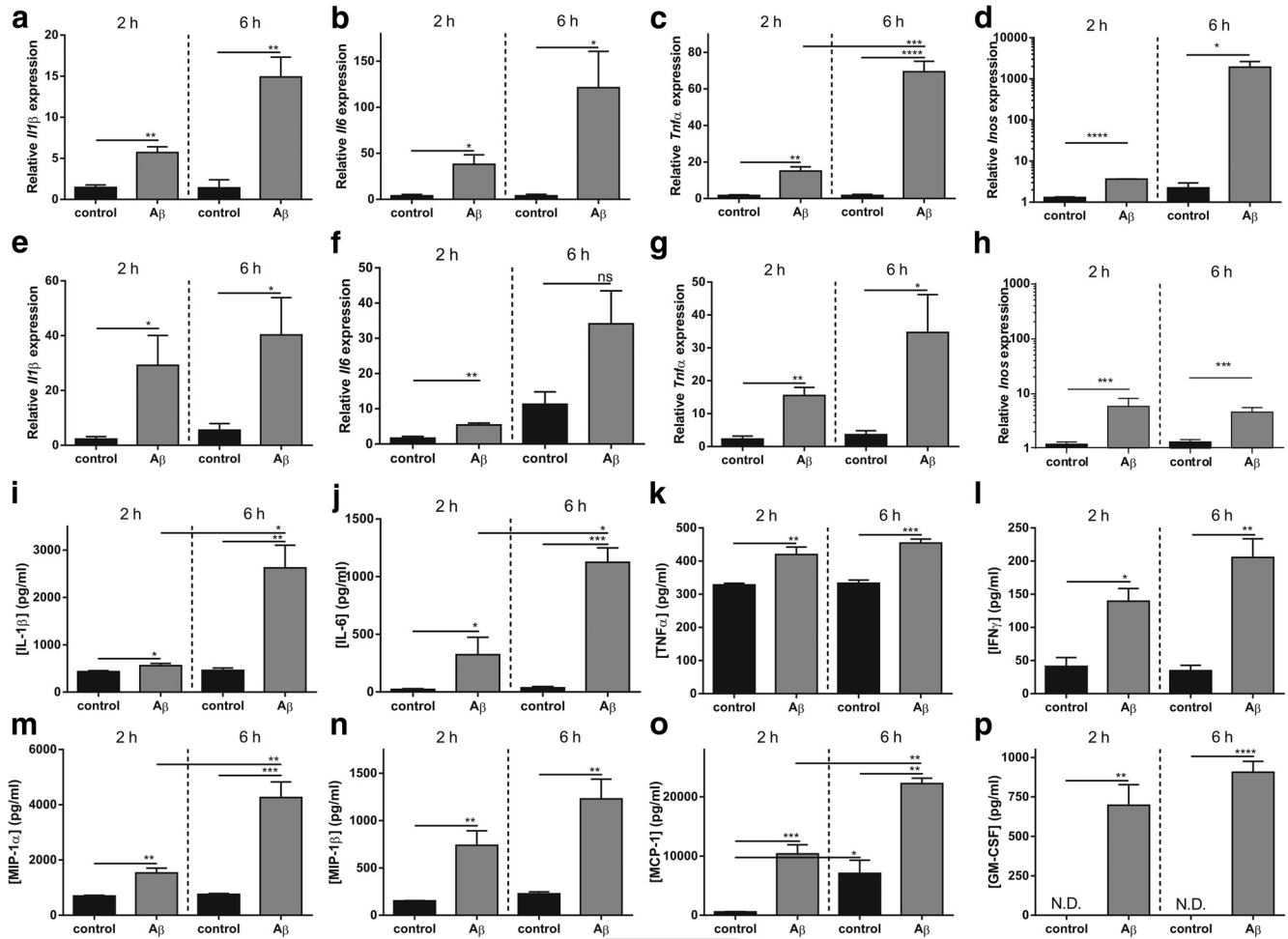
Intracerebroventricular injection of A $\beta$ 1–42 oligomers induces morphological changes in the CP and loss of BCSFB integrity

The CP undergoes morphological alterations in Alzheimer's disease (Marques et al., 2013). Therefore, we injected C57BL/6 mice intracerebroventricularly with oligomerized A $\beta$ 1–42 and studied CPE morphology 6 h later using SBF-SEM. This technology uses a fully automated microtome installed in the SEM chamber to produce serial sections that makes 3D reconstructions of SEM samples (Denk and Horstmann, 2004) and this revealed that A $\beta$ 1–42 induces loss of the typical cuboidal structure of CPE cells (Fig. 3; Movies 1 and 2). To better visualize the differences in our SBF-SEM images, we performed 3D modeling of CPE cells to highlight the morphological differences. CPE cell outlines were drawn manually using IMOD software for ~200 sections per sample. These sections were then merged to generate a 3D model of the cell shape. CPE cells retained typical cuboidal structure after scrambled peptide injection. In contrast, several cells in A $\beta$ 1–42 oligomer-injected mice lost their cuboidal shape and the overall cell volume was reduced compared with the scrambled

F1

F2

F3, Mv1-Mv2



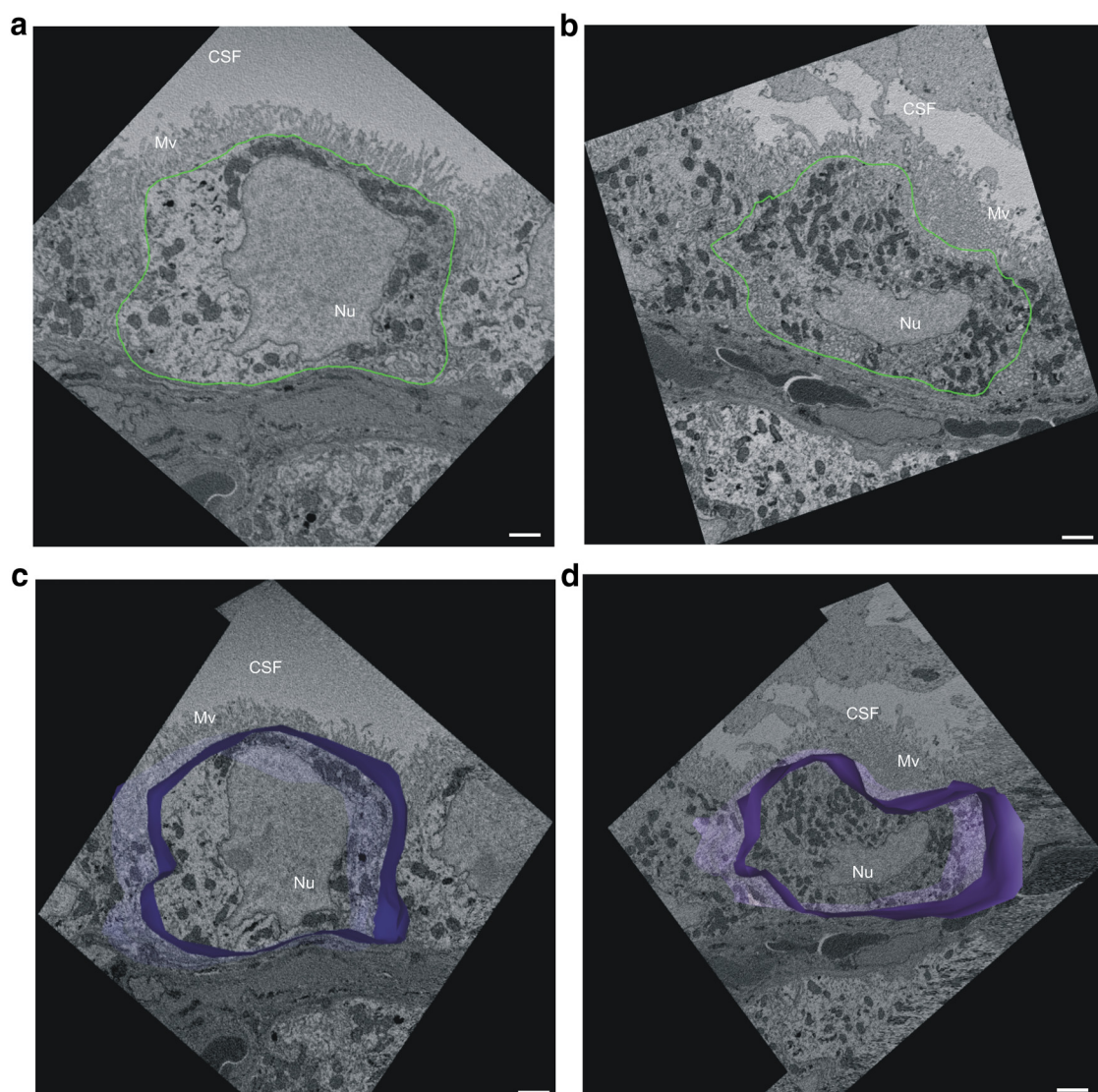
**Figure 2.** Cytokine and chemokine analyses of CP, hippocampus, and CSF after intracerebroventricular injection of A $\beta$ 1–42 oligomers. **a–d**, mRNA expression analysis of *Il1 $\beta$*  (a), *Il6* (b), *Tnf* (c), and *Inos* (d) in CP after intracerebroventricular injection of A $\beta$ 1–42 (gray) in C57BL/6 mice compared with control CP samples (black) ( $n = 3-4$ ). **e–h**, mRNA expression analysis of *Il1 $\beta$*  (e), *Il6* (f), *Tnf* (g), and *Inos* (h) in the hippocampus after intracerebroventricular injection of A $\beta$ 1–42 (gray) in C57BL/6 mice compared with control samples (black) ( $n = 3-4$ ). **i–p**, Levels of cytokines (IL-1 $\beta$ , IL-6, TNF $\alpha$ , and IFN $\gamma$ ) and chemokines (MIP-1 $\alpha$ , MIP-1 $\beta$ , MCP-1, and GM-CSF) in CSF isolated from C57BL/6 mice injected intracerebroventricularly with scrambled peptide (black) or A $\beta$ 1–42 (gray) ( $n = 4$ ).

injected CPE cells. In addition, CPE cells in A $\beta$ 1–42 oligomer-injected mice displayed reduced nucleus size (Fig. 3c,d). Next, we analyzed whether the severe morphological changes induced by A $\beta$ 1–42 oligomers affect BCSFB integrity. To compare blood–CSF permeability in control mice and in mice injected intracerebroventricularly with A $\beta$ 1–42 oligomers, we injected 4 kDa FITC-dextran intravenously 1 h before CSF isolation and measured fluorescence in CSF. The results revealed that A $\beta$ 1–42 oligomers induce permeability of the BCSFB 6 h after intracerebroventricular injection, demonstrating that oligomerized A $\beta$ 1–42 disturbs CPE barrier functionality (Fig. 4a).

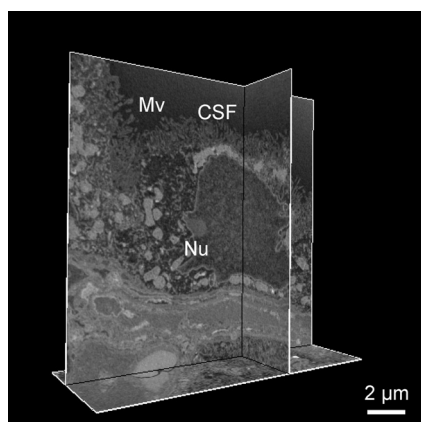
Subsequently, we used qPCR, Western blotting, and immunostaining to determine whether the alterations in BCSFB integrity induced by A $\beta$ 1–42 oligomers were correlated with changes in junctional proteins in the CP. Gene expression analysis revealed that several tight junction genes were affected by intracerebroventricular injection of A $\beta$ 1–42 oligomers. As shown in Figure 4, b and c, expression of Claudin-5 (*Cldn5*) and Occludin (*Ocln*) was significantly downregulated in response to A $\beta$ 1–42 oligomers. Notably, *Ocln* was significantly downregulated at both time points tested, whereas *Cldn5* was significantly downregulated only 6 h after A $\beta$ 1–42 oligomer injection. Zona occludens-1 (*Zo1*) and Claudin-1 (*Cldn1*) were significantly downregulated

2 h after A $\beta$ 1–42 oligomer injection (Fig. 4e,f). In contrast, *Zo3* showed no change in gene expression at that time point (Fig. 4g). Gene expression of the adherens junctional genes E-cadherin (*Ecdh*) and N-cadherin (*Ncdh*) was unaffected at both time points (Fig. 4h,i). OCLN, the tight junction protein that was most severely affected 6 h after intracerebroventricular injection, was also analyzed by Western blot. CP tissue was isolated from control mice and from mice injected intracerebroventricularly with A $\beta$ 1–42 oligomers and pooled samples were analyzed by SDS-PAGE. Immunodetection revealed that OCLN protein levels are downregulated by intracerebroventricular injection of A $\beta$ 1–42 oligomers (Fig. 4d). This was also studied by fluorescent immunostaining. Confocal imaging revealed that, in control mice, OCLN was enriched at the apical side of the CPE cells (Fig. 4j,k, white arrowheads). In agreement with the Western blot analysis, the OCLN signal was reduced after A $\beta$ 1–42 oligomer injection (Fig. 4l,m, white arrowheads). We also analyzed the adherens junction protein ECDH. Like OCLN, this protein was enriched at the apical side of CPE cells, but it was unaffected by intracerebroventricular injection of A $\beta$ 1–42 oligomers (Fig. 4n,o). Our data show that tight junction proteins are affected at the mRNA and/or protein level upon intracere-

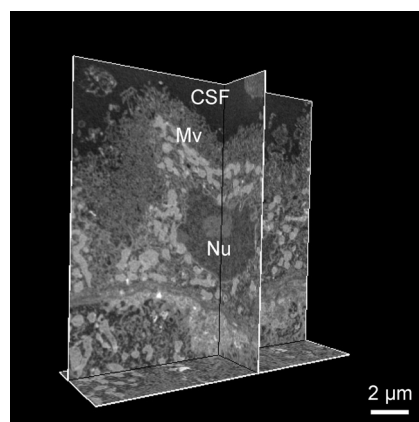




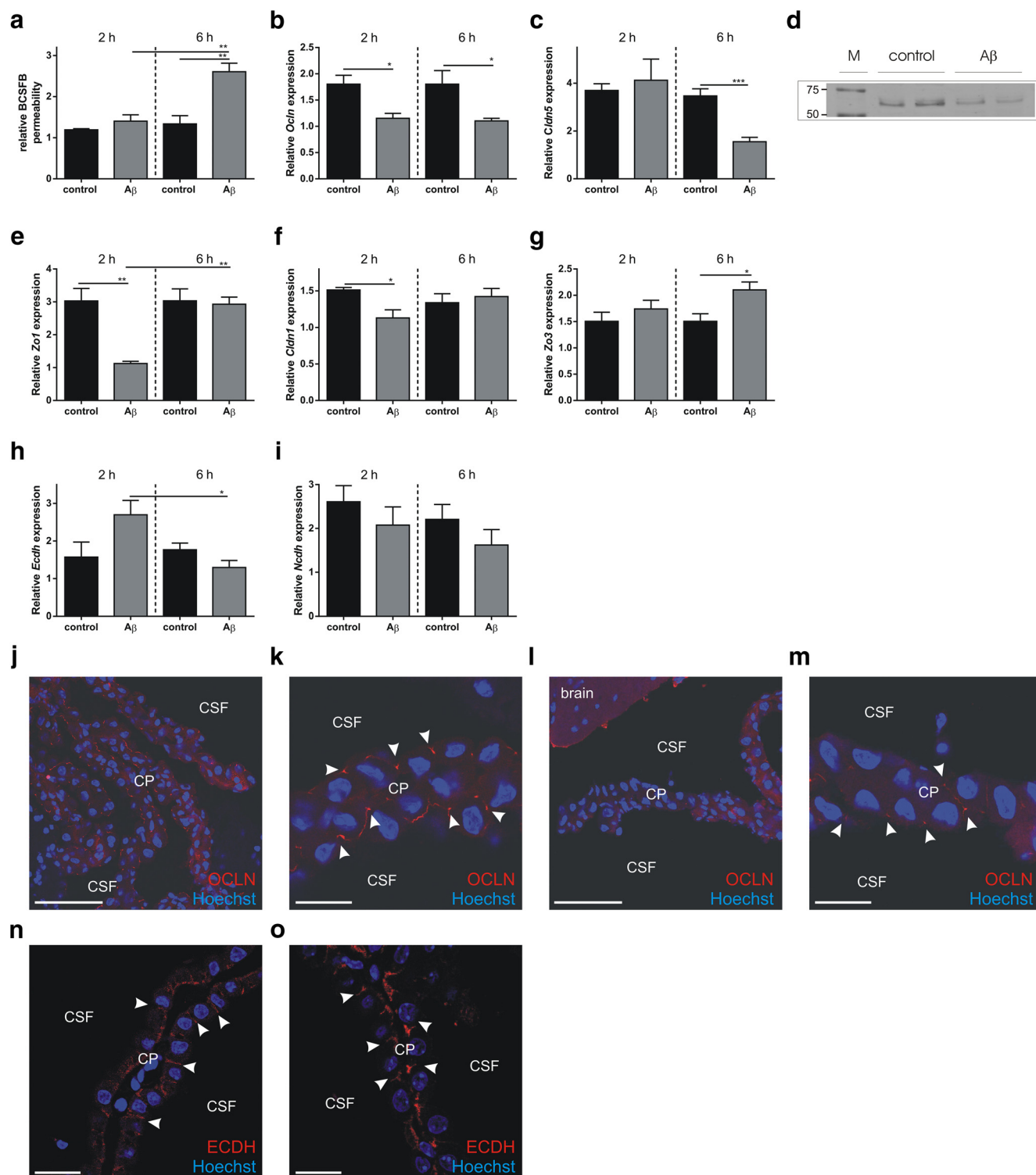
**Figure 3.** CP morphology analysis by SFB-SEM. *a, b*, Representative SFB-SEM images of CP cells of C57BL/6 mice injected intracerebroventricularly with scrambled (*a*) or A $\beta$ 1–42 oligomers (*b*). Cell shape is outlined in green. *c, d*, 3D modeling (blue) based on merging ~200 sections of CPE cells from scrambled (*c*) and A $\beta$ 1–42 oligomer (*d*) injected mice. Only cell shape was considered; basolateral labyrinth and microvilli were neglected while generating the 3D modeling. Mv, Microvilli; Nu, nucleus. Scale bar, 2  $\mu$ m.



**Movie 1.** Morphology of CPE of mice after injection of scrambled peptide into the cerebral ventricles. Shown are representative 3D reconstructions of CPE samples from C57BL/6 mice by SFB-SEM 6 h after injection of scrambled peptide in the cerebral ventricles.



**Movie 2.** Morphology of the CPE of mice injected in the cerebral ventricles with A $\beta$ 1–42 peptide. Shown are representative 3D reconstructions of CPE samples from C57BL/6 mice by SFB-SEM 6 h after injection of A $\beta$ 1–42 peptide in the cerebral ventricles.

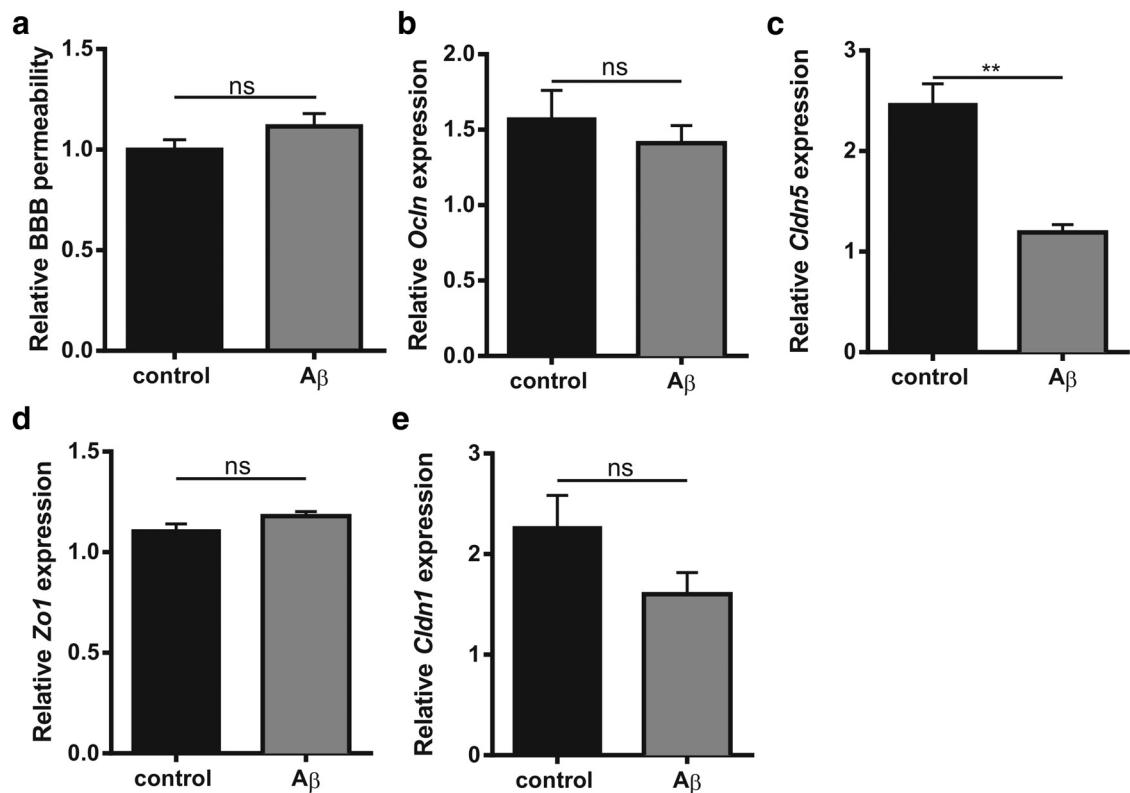


**Figure 4.** Analysis of  $A\beta$ 1–42-induced disruption of BCSFB integrity. **a**, Relative BCSFB permeability 2 and 6 h after intracerebroventricular injection of  $A\beta$ 1–42 in the cerebral ventricles (gray) compared with control mice (black) ( $n = 4$ ). **b**, **c**, *Ocln* (**b**) and *Cldn5* gene (**c**) expression in CP tissue of C57BL/6 mice injected intracerebroventricularly with scrambled (black) or  $A\beta$ 1–42 (gray) peptide ( $n = 4$ ). **d**, Western blot analysis of Occludin in CP tissue from C57BL/6 control mice and mice injected intracerebroventricularly with  $A\beta$ 1–42 peptide. **e–i**, *Zo1* (**e**), *Cldn1* (**f**), *Zo3* (**g**), *Ecdh* (**h**), and *Ncdh* (**i**) gene expression in CP tissue of C57BL/6 mice injected in the cerebral ventricles with  $A\beta$ 1–42 (gray) or scrambled peptide (black) ( $n = 4$ ). **j–m**, Representative confocal images of C57BL/6 control mice (**j**, **k**) and mice injected intracerebroventricularly with  $A\beta$ 1–42 (**l**, **m**) peptide for 6 h (red, Occludin; blue, Hoechst). The arrowheads point to the apically located tight junctions. Scale bars: **e**, **g**, 50  $\mu$ m; **f**, **h**, 10  $\mu$ m. **n**, **o**, Representative confocal images of control (**n**) and  $A\beta$ 1–42 (**o**) peptide injected intracerebroventricularly in C57BL/6 mice stained for E-cadherin (ECDH; red) and Hoechst (blue). The arrowheads point to adherens junctions.

broventricular  $A\beta$ 1–42 oligomer injection, which might explain the observed BCSFB leakage.

In parallel, we also studied the effect of  $A\beta$ 1–42 oligomers on BBB integrity. As shown in Figure 5*a*, we did not observe a sig-

nificant increase in BBB leakage 6 h after  $A\beta$ 1–42 oligomer administration. In addition, also *Ocln*, *Cldn1*, and *Zo1* gene expression was not affected in the hippocampus upon intracerebroventricular injection of  $A\beta$ 1–42 oligomers, whereas *Cldn5*



**Figure 5.** Analysis of the effect of A $\beta$ 1–42 oligomers on BBB integrity. **a**, Relative permeability of the BBB 6 h after intracerebroventricular injection of A $\beta$ 1–42 oligomers (gray) compared with scrambled peptide injected mice (black) ( $n = 10–13$ ). **b–e**, Relative gene expression of tight junction proteins *Occludin* (**b**), *Claudin-5* (**c**), *Zona occludens-1* (**d**), and *Claudin-1* (**e**) in the hippocampus 6 h after intracerebroventricular injection of scrambled peptide (black) or A $\beta$ 1–42 oligomers (gray) ( $n = 5$ ).

did show a significant decrease in gene expression compared with scrambled peptide control (Fig. 5*b–e*). Clearly, intracerebroventricular injection of A $\beta$ 1–42 oligomers only modestly affects the BBB 6 h after the injection.

**A $\beta$ 1–42 oligomer-induced BCSFB disruption is linked to increased MMP expression and activity**

Previous studies have shown altered expression of MMPs in Alzheimer’s disease patients (Wang et al., 2014) and MMPs have been implicated in BCSFB disruption in response to TNF (Zeni et al., 2007) and in several inflammatory conditions (Vandenbroucke and Libert, 2014) such as sepsis (Vandenbroucke et al., 2012) and stroke (Batra et al., 2010). To determine whether MMPs play a role in the A $\beta$ 1–42 oligomer-induced BCSFB leakage, we analyzed MMP gene expression in the CP and MMP activity in the CSF. Figure 6*a* shows the fold changes in gene expression of *Mmp1a*, *Mmp2*, *Mmp3*, *Mmp8*, *Mmp9*, *Mmp13*, and *Mmp14* 2 h (black) and 6 h (gray) after intracerebroventricular injection of A $\beta$ 1–42 oligomers. *Mmp3* was significantly up-regulated 2 h after intracerebroventricular injection and *Mmp3*, *Mmp8*, *Mmp9*, and *Mmp13* at 6 h after injection. This was associated with an increase in MMP activity in the CSF 6 h after A $\beta$ 1–42 oligomer injection (Fig. 6*b*). Next, we coinjected A $\beta$ 1–42 oligomers intracerebroventricularly with the broad-spectrum MMP inhibitor GM6001. The results show that inhibition of MMP activity prevented the A $\beta$ 1–42 oligomer-induced BCSFB disruption. Indeed, comparison of BCSFB integrity of mice injected with A $\beta$ 1–42 oligomers alone or combined with MMP inhibitor revealed that MMP inhibition prevents oligomerized A $\beta$ 1–42 from inducing BCSFB leakage (Fig. 6*c*). As expected, this was associated with a decrease in MMP activity in

the CSF (Fig. 6*b*). Clearly, inhibiting MMP activity diminishes the detrimental effects of A $\beta$ 1–42 oligomers. Based on the gene expression analysis displayed in Figure 6*a*, we also analyzed MMP3 protein level in the CSF. Western blot analysis clearly showed an increase in secreted MMP3 (Fig. 6*d*). Finally, we injected A $\beta$ 1–42 oligomers intracerebroventricularly in wild-type and MMP3-deficient (MMP3<sup>−/−</sup>) mice to confirm the suspected role of MMPs, especially MMP3. As shown in Figure 6*e*, absence of MMP3 significantly prevented BCSFB leakage induced by A $\beta$ 1–42 oligomers. In agreement with this, A $\beta$ 1–42 oligomers did not affect the morphology of the CPE cells in MMP3<sup>−/−</sup> mice (Fig. 7).

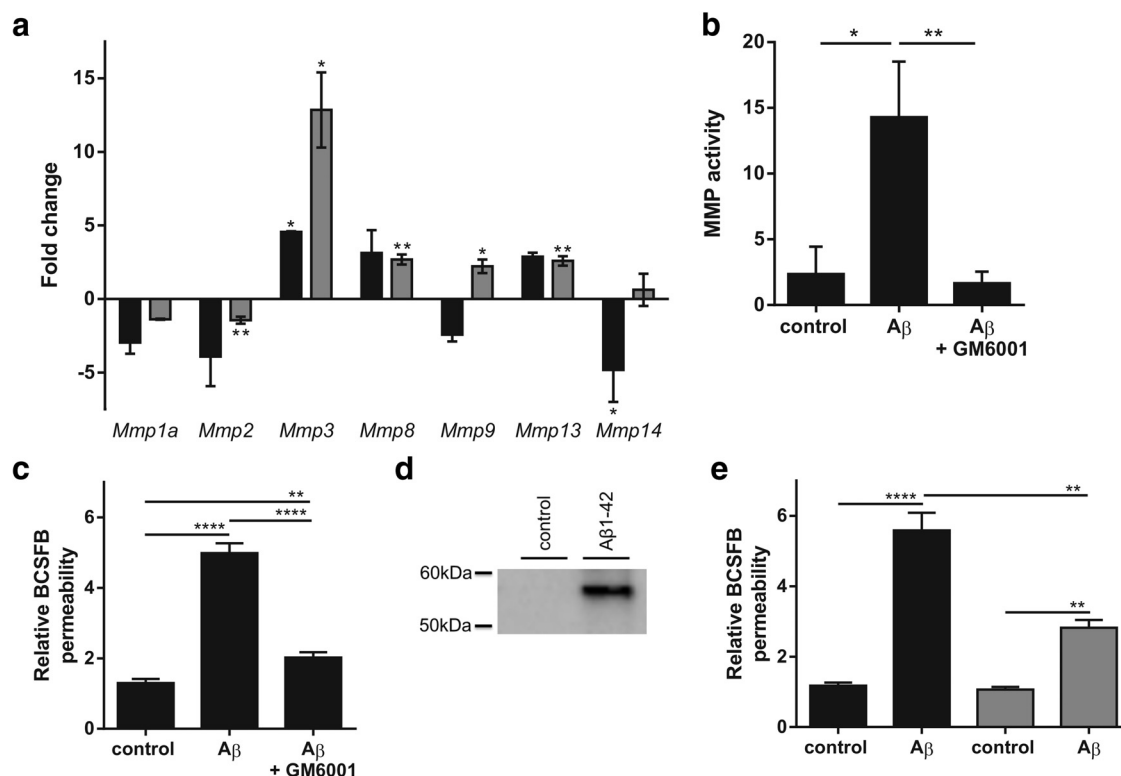
**Discussion**

Neuroinflammation occurs very early on in disease progression and in several brain regions affected by Alzheimer’s disease. Abnormal molecules such as A $\beta$ 1–42, hyperphosphorylated tau protein, and damaged neurons induce the activation of cytokines, complement proteins, and chemokines, as well as other acute phase pathways, eventually leading to inflammation (Akiyama et al., 2000). In a 2004 human study, increased levels of proinflammatory cytokines were found in plasma before the clinical onset of the disease (Engelhart et al., 2004). In addition, cell culture studies and animal models of Alzheimer’s disease demonstrated that inflammation precedes the appearance of pathological hallmarks of Alzheimer’s disease such as senile plaques and neurofibrillary tangles (Floden et al., 2005; Wright et al., 2013). Some evidence indicates that soluble A $\beta$  oligomers, an intermediate between monomeric A $\beta$  and insoluble amyloid plaques, are an important mediator in the inflammation and pathology of Alzheimer’s disease. Although the concentration of

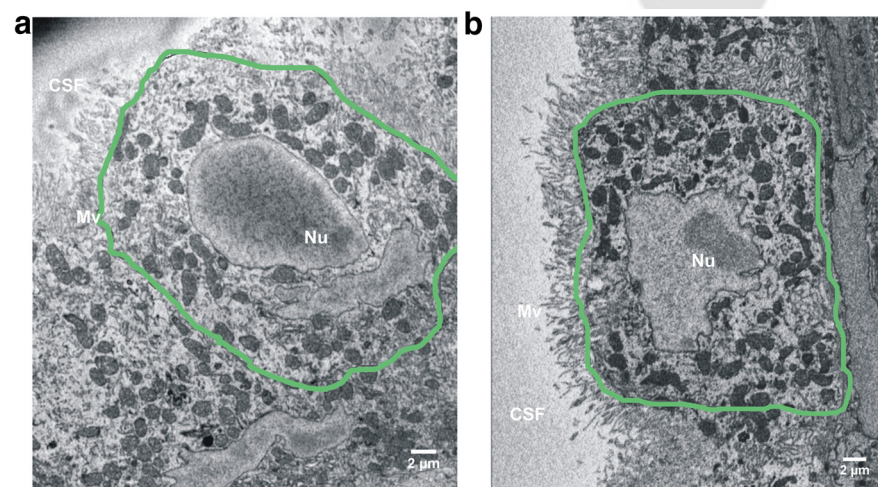
F6

F7





**Figure 6.** Analysis of the role of MMPs in  $A\beta$ 1–42-induced disruption of BCSFB permeability. **a**, Fold change in *Mmp* gene expression in the CP 2 h (black) and 6 h (gray) after intracerebroventricular injection of  $A\beta$ 1–42 in the cerebral ventricles compared with control samples ( $n = 3–4$ ). **b**, Total MMP activity in CSF of C57BL/6 mice 6 h after intracerebroventricular injection of scrambled peptide (control),  $A\beta$ 1–42 oligomers ( $A\beta$ ), or  $A\beta$ 1–42 oligomers together with GM6001 ( $A\beta$  + GM6001) ( $n = 6$ ). **c**, BCSFB permeability of C57BL/6 mice 6 h after intracerebroventricular injection of  $A\beta$ 1–42 oligomers alone ( $A\beta$ ) or combined with the MMP inhibitor GM6001 ( $A\beta$  + GM6001) compared with scrambled peptide injected mice (control) ( $n = 6–7$ ). **d**, Western blot analysis of MMP3 protein levels in CSF of control and  $A\beta$ 1–42 oligomer intracerebroventricularly injected mice. **e**, Relative BCSFB permeability in wild-type (black) and  $MMP3^{-/-}$  (gray) C57BL/6 mice 6 h after intracerebroventricular injection of scrambled control or  $A\beta$ 1–42 oligomers ( $n = 4–10$ ).



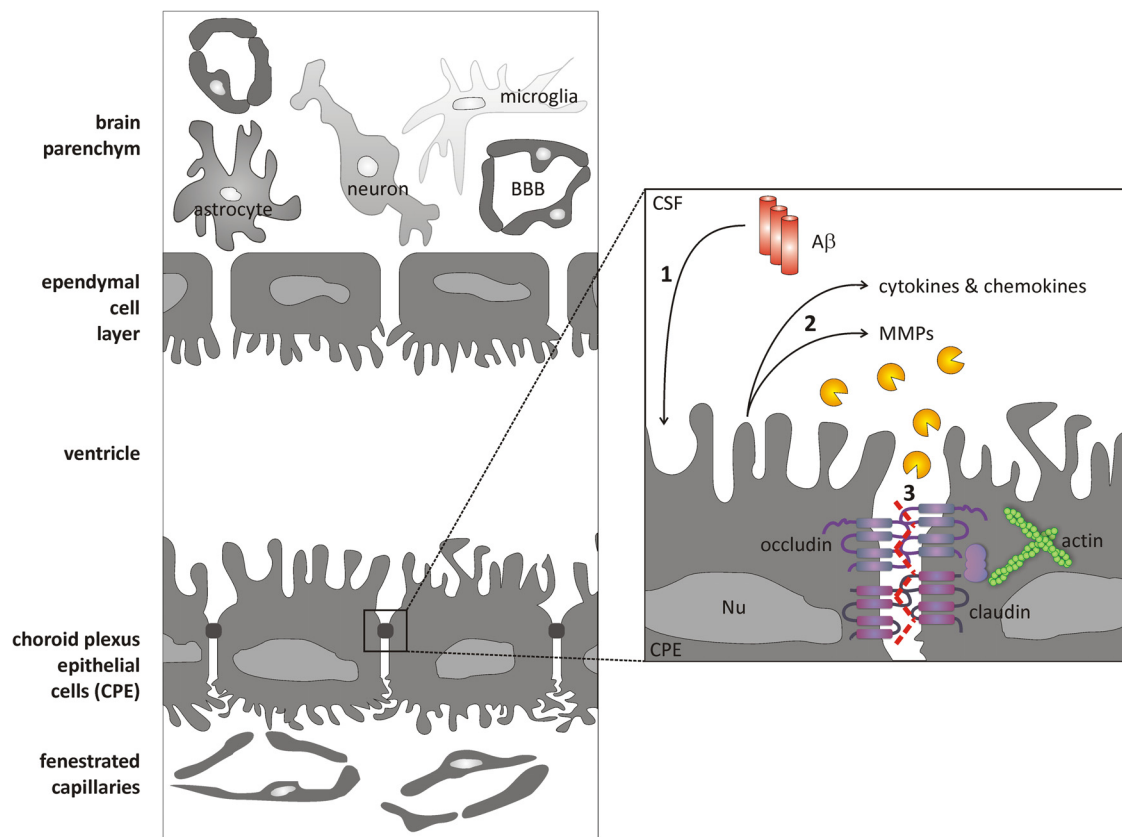
**Figure 7.** CP morphology analysis by SFB-SEM of  $MMP3^{-/-}$  mice. **a**, **b**, Representative SFB-SEM images of the CPE cells of scrambled peptide (**a**) and  $A\beta$ 1–42 oligomer (**b**) intracerebroventricularly injected  $MMP3^{-/-}$  mice. Cell shape is represented in green. Basolateral labyrinth and microvilli were neglected while drawing the cell outline. Mv, Microvilli; Nu, nucleus.

soluble  $A\beta$ 1–42 was not uniform in different studies (Nitta et al., 1997; Kong et al., 2005; Marco and Skaper, 2006; Brouillette et al., 2012; Cetin et al., 2013), in most,  $A\beta$ 1–42 was able to initiate a cascade of the events that recapitulate the key pathological hallmarks of Alzheimer's disease. *In vitro* incubation of macrophages, microglia, astrocytes, and oligodendrocytes with soluble  $A\beta$  induced inflammation (Johnstone et al., 1999; Lee et al., 2002;

Smits et al., 2002). *In vivo*, microvessels forming the BBB isolated from the brains of Alzheimer's disease patients showed markedly increased levels of IL-1 $\beta$ , IL-6, and TNF compared with control subjects (Grammas and Ovase, 2001; Tripathy et al., 2007).

Here, we made use of  $A\beta$ 1–42 oligomers prepared as described previously, which are known to induce neuronal cell loss, tau hyperphosphorylation, and impairment of hippocampus-dependent memory (Brouillette et al., 2012; Ledo et al., 2013; Nery et al., 2014). Injected  $A\beta$ 1–42 levels were ~fourfold compared with levels measured in transgenic APP/PS1 mice (Maia et al., 2013). In agreement with previous studies, intracerebroventricular injection of  $A\beta$ 1–42 oligomers induced brain inflammation, as reflected by increased gene expression of *Il6*, *Il1 $\beta$* , and *TNF* in the hippocampus, the area of the brain crucial for memory and among

the first regions to be affected in Alzheimer's disease. Using the same mouse model as in our study, Kuperstein et al. (2010) found inhibition of new memory formation 24 h after  $A\beta$ 1–42 oligomer injection. Because elevated cytokine levels in hippocampus are able to induce memory impairment in rats (Czerniawski and Guzowski, 2014), it could be speculated that elevated proinflam-



**Figure 8.** Schematic overview of direct effects of A $\beta$ 1–42 oligomers on the BCSFB. A monolayer of CPE cells, tightly connected by tight junctions, restricts entrance of molecules from the fenestrated capillaries into the CSF. Injection of oligomerized A $\beta$ 1–42 into the cerebral ventricles of mice (1) leads to secretion of MMPs, cytokines, and chemokines from CPE cells into the CSF (2), and this induces disruption of tight junctions (3), eventually resulting in BCSFB leakage.

matory cytokine levels contribute to memory deterioration in this model of Alzheimer's disease.

Several studies have shown that Alzheimer's disease is associated with morphological and functional changes at the BCSFB (Serot et al., 2000; Serot et al., 2003; Emerich et al., 2005; Krzyzanowska and Carro, 2012; Serot et al., 2012; Marques et al., 2013; Spector and Johanson, 2013). However, it is not fully understood how the integrity of the BCSFB is affected in Alzheimer's disease and whether the CP is involved in triggering and/or enhancing the inflammatory response to A $\beta$ . Therefore, we studied the effect of intracerebroventricular injection of A $\beta$ 1–42 oligomers on the BCSFB. The BCSFB is formed by a single continuous layer of modified cuboidal epithelium, the CPE cells, which are attached on their basal sides to the basal lamina (Wolburg and Paulus, 2010). CPE cells are large and have a centrally positioned spherical nucleus, copious cytoplasm, and plenty of apically positioned microvilli. In addition to their barrier function, CPE cells are responsible for the production and secretion of CSF, as well as other biologically active components involved in nutrition, endocrine function, and clearance of the brain areas (Johanson et al., 2008). Here, we show that the CP responds to intracerebroventricular injection of soluble A $\beta$ 1–42 oligomers by increasing its expression of the proinflammatory cytokines *Il1 $\beta$* , *Il6*, *Tnfa*, and *Inos*. This was associated with significantly elevated levels of cytokines and chemokines in the CSF. One of the Alzheimer's disease theories proposes that monomeric and oligomeric A $\beta$  activate microglia, which respond by secreting proinflammatory cytokines, leading to the spread of inflammation, cell death, and overall neurodegeneration (Maezawa et al., 2011; Solito and Sas-

tre, 2012). However, despite the observed increased mRNA levels of proinflammatory cytokines in the hippocampus upon intracerebroventricular injection of A $\beta$ 1–42 oligomers, we did not observe significant microglial activation. This indicates that the inflammatory signal generated in the CPE in response to soluble A $\beta$ 1–42 oligomers might occur before microglia activation during the development of Alzheimer's disease. However, we do not exclude that the injected A $\beta$ 1–42 oligomers also affect other cell types, as noted in the past by other research groups (Kopec and Carroll, 1998; von Bernhardi and Eugénin, 2004).

Some of the reported Alzheimer's disease-associated changes involve CPE cell atrophy, accumulation of Biondi tangles and lipofuscin deposits, irregular basement thickening, decreased CSF production and secretion, and diminished capacity for clearance of unwanted molecules from CSF (Emerich et al., 2005). In our study, we observed that a single intracerebroventricular injection of A $\beta$ 1–42 oligomers affected the cuboidal morphology of CPE cells, suggesting that some of the observed morphological changes at the CPE in Alzheimer's disease occur early during disease development.

The ability of CPE cells to restrict paracellular passage of molecules depends on the presence of junctional complexes at their apical sides (Redzic, 2011). Several studies have shown a link between increased cytokine levels and decreased tight junction expression at the BBB and BCSFB (Minagar and Alexander, 2003; Barichello et al., 2011; Chai et al., 2014). Diminished BBB integrity, which leads to an increase in permeability and a decrease in tight junction expression, was found in both mouse and human Alzheimer's disease studies (Romanitan et al., 2010; Biron et al.,

2011; Carrano et al., 2011; Hartz et al., 2012). In agreement with the  $\text{A}\beta$ 1–42 oligomer-induced cytokine increase in the CSF, we observed that a single intracerebroventricular injection of  $\text{A}\beta$ 1–42 oligomers affected the gene and protein expression levels of tight junctions. Furthermore, we observed increased leakage of fluorescently labeled dextran from blood into the CSF, indicating that  $\text{A}\beta$ 1–42 injection leads to loss of BCSFB integrity. In contrast, we did not observe loss of BBB integrity 6 h after intracerebroventricular injection of  $\text{A}\beta$ 1–42 oligomers.

MMPs have been recognized as important players in neurodegenerative diseases (Rosenberg, 2009). The large family of MMPs includes collagenases, gelatinases, stromelysins, and membrane-type MMPs, which are expressed by endothelial, epithelial, and inflammatory cells at the different brain barriers in response to inflammation (Vandenbroucke and Libert, 2014). The involvement of MMPs in BCSFB breakdown has been observed in various disease models (Zeni et al., 2007; Batra et al., 2010; Vandenbroucke et al., 2012). So far, there are no data implicating MMPs in the loss of BCSFB integrity during Alzheimer's disease. In our acute model of Alzheimer's disease, we observed increased mRNA levels of several MMPs associated with increased MMP activity in the CSF. The  $\text{A}\beta$ 1–42 oligomer-induced increase was most prominent for *Mmp3*. Importantly, the broad-spectrum MMP inhibitor GM6001 prevented the detrimental effects of  $\text{A}\beta$ 1–42 oligomer injection on the BCSFB, suggesting the involvement of MMPs in  $\text{A}\beta$ 1–42 oligomer-induced BCSFB breakdown. Although we did not show this in our study, the observed BCSFB dysfunction might be due to direct cleavage of tight junction proteins, as was shown previously for several MMPs (Nava et al., 2013), or it might also be caused by cytokine activation or cleavage of extracellular matrix components, as we have shown previously (Vandenbroucke et al., 2012; Vandenbroucke et al., 2013).

MMP3 is believed to be a key player in inflammation and is secreted by CPE cells (Thouvenot et al., 2006). Indeed, we were able to detect secreted MMP3 in the CSF of  $\text{A}\beta$ 1–42 oligomer-injected mice. Interestingly, several recent studies propose MMP3 as a potential biomarker for Alzheimer's disease because it is significantly upregulated in the brain, CSF, and plasma of Alzheimer's disease patients (Yoshiyama et al., 2000; Horstmann et al., 2010; Hanzel et al., 2014; Kauwe et al., 2014; Mroczko et al., 2014). A recent study in a mouse model of spinal cord injury showed that MMP3-deficient mice display reduced disruption of the blood–spinal cord barrier (Lee et al., 2014). To further study the role of MMP3, we injected MMP3-deficient mice intracerebroventricularly with  $\text{A}\beta$ 1–42 oligomer and found that the absence of MMP3 strongly reduced the  $\text{A}\beta$ 1–42 oligomer-induced BCSFB leakage. Moreover, *Mmp8*, *Mmp9*, and *Mmp13* mRNA levels were elevated in the CP upon intracerebroventricular injection of  $\text{A}\beta$ 1–42 oligomer, so these MMPs might also play a role in the observed loss of BCSFB integrity. In previous work, we showed that MMP8 contributes to inflammation-induced BCSFB leakage (Vandenbroucke et al., 2012) and MMP8 (Schubert-Unkmeir et al., 2009), MMP9 (Chiu and Lai, 2013), and MMP13 (Vandenbroucke et al., 2013) have been linked to tight junction dysregulation. Previously, MMP9 was shown to be involved in BBB breakdown mediated by ApoE4 (Halliday et al., 2013). Altogether, our data provide solid evidence for a detrimental role of MMPs in  $\text{A}\beta$ 1–42 oligomer-induced BCSFB disruption.

As schematically depicted in Figure 8, our data indicate that  $\text{A}\beta$ 1–42 oligomers, via CPE-derived cytokine and MMP secretion, might induce BCSFB barrier breakdown early during the

development of Alzheimer's disease, thereby contributing to enhancement of neuroinflammation. Indeed, loss of brain barrier integrity might further aggravate neuroinflammation by entrance of undesirable molecules into the brain. It has been suggested that BBB dysfunction might contribute to Alzheimer's disease (Erickson and Banks, 2013). Based on our results, BCSFB dysfunction might be an even earlier event during disease development, so further research is needed to study whether restoring BCSFB integrity might reduce neuroinflammation during Alzheimer's disease.

In conclusion, our study shows that the presence of  $\text{A}\beta$ 1–42 oligomers in the CSF induces disruption of the BCSFB via the production of proinflammatory cytokines and MMPs, which is linked to loss of tight junction functionality.

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